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SEQUENCE ANALYSIS

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EMBOSS-Align Results

Needle Results	
Matrix	DNAfull
Open gap penalty	10.0
Gap extension penalty	0.5
Needle output	needle-20060804-00212328575043.output

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EMBOSS-Align Results

Needle Results	
Matrix	Blosum62
Open gap penalty	10.0
Gap extension penalty	0.5
Needle output	needle-20060804-00264023727706.output

```

#####
# Program: needle
# Rundate: Fri Aug 04 00:26:41 2006
# Align_format: srspair
# Report_file: /ebi/extserv/old-work/needle-20060804-00264023727706.output
#####

=====
#
# Aligned_sequences: 2
# 1: EMBOSS_001
# 2: EMBOSS_001
# Matrix: EBLOSUM62
# Gap_penalty: 10.0
# Extend_penalty: 0.5
#
# Length: 351
# Identity: 351/351 (100.0%)
# Similarity: 351/351 (100.0%)
# Gaps: 0/351 ( 0.0%)
# Score: 1821.0
#
#
=====

EMBOSS_001      1 MGDPERPEAAGLDQDERSSSDTNESEIKSNEEPLLRKSSRRFVIFPIQYP
                  |||||||Q|||||Q|||||Q|||||Q|||||Q|||||Q|||||Q|||||Q|||||Q
EMBOSS_001      1 MGDPERPEAAGLDQDERSSSDTNESEIKSNEEPLLRKSSRRFVIFPIQYP
                  |||||||Q|||||Q|||||Q|||||Q|||||Q|||||Q|||||Q|||||Q|||||Q
EMBOSS_001      51 DIWKMYKQAQASFWTAEVVDLSKDLPHWNKLKADEKYFISHILAFFAASD
                  |||||||Q|||||Q|||||Q|||||Q|||||Q|||||Q|||||Q|||||Q|||||Q
EMBOSS_001      51 DIWKMYKQAQASFWTAEVVDLSKDLPHWNKLKADEKYFISHILAFFAASD
                  |||||||Q|||||Q|||||Q|||||Q|||||Q|||||Q|||||Q|||||Q|||||Q
EMBOSS_001      101 GIVNENLVERFSQEVQVPEARCFYGFQILIENVHSEMSLLIDTYIRDPK
                  |||||||Q|||||Q|||||Q|||||Q|||||Q|||||Q|||||Q|||||Q|||||Q
EMBOSS_001      101 GIVNENLVERFSQEVQVPEARCFYGFQILIENVHSEMSLLIDTYIRDPK
                  |||||||Q|||||Q|||||Q|||||Q|||||Q|||||Q|||||Q|||||Q|||||Q
EMBOSS_001      151 KREFLFNIAETMPYVKKKADWALRWIADRKFSTGERVVAFAAVEGVFFSG
                  |||||||Q|||||Q|||||Q|||||Q|||||Q|||||Q|||||Q|||||Q|||||Q
EMBOSS_001      151 KREFLFNIAETMPYVKKKADWALRWIADRKFSTGERVVAFAAVEGVFFSG
                  |||||||Q|||||Q|||||Q|||||Q|||||Q|||||Q|||||Q|||||Q|||||Q
EMBOSS_001      201 SFAAIFWLKKRGLMPGLTFSNELISRDEGLHCDFACLMFQYLVNKPSEER
                  |||||||Q|||||Q|||||Q|||||Q|||||Q|||||Q|||||Q|||||Q|||||Q

```

EMBOSS_001	201	SFAAIFWLKKRGLMPGLTFSNELISRDEGLHCDFACLMFQYLVNKPSEER	250
EMBOSS_001	251	VREIIVDAVKIEQEFLTEALPVGLIGMNCILMKQYIEFVADRLVELGFS	300
EMBOSS_001	251	VREIIVDAVKIEQEFLTEALPVGLIGMNCILMKQYIEFVADRLVELGFS	300
EMBOSS_001	301	KVFQAENPFDGMENISLEGKTNFFEKRVSEYQRFAVMAETTDNVFTLDAD	350
EMBOSS_001	301	KVFQAENPFDGMENISLEGKTNFFEKRVSEYQRFAVMAETTDNVFTLDAD	350
EMBOSS_001	351	F 351	
EMBOSS_001	351	F 351	

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